

A photograph of a busy city street at night, filled with cars and streetlights. The scene is illuminated by warm yellow streetlights and the headlights of the vehicles. In the background, there are buildings and a large digital billboard. The overall atmosphere is one of a bustling urban environment.

# IdentiPROT

**комплексное решение для потока протеемных данных  
по принципу «все включено»**

The image shows a browser window with several tabs. The active tab is titled "Welcome to Ide...". The address bar shows the URL "identiprot.theorchromo.ru/logout/". The page content includes a large blue header with the text "IdentiPROT" and navigation links for "ABOUT", "LOG IN", and "CONTACTS". A main heading reads "Welcome to the home of IdentiPROT software!". Below this, a paragraph states: "If you have problems logging in, any questions or comments, please [contact](#) our administrators to create account." On the right side, there is a dark blue box with the text "WELCOME The single entry point for your proteomic data mining". Below this box are two input fields labeled "Email" and "Password", followed by a large blue "Log in" button.

**Вход через Интернет из клиентского места, включая мобильные устройства**

New Search

identiprot.theorchromo.ru/data/

**load parameters**

send email notification

use auto optimization

enzyme: [add custom cleavage rule](#) trypsin

number of missed cleavages 2

precursor accuracy unit ppm

precursor accuracy left 10

precursor accuracy right 10

precursor isotope mass error 0

product accuracy, Da 0.1

FDR 1.0

FDR type psm

minimum charge 2

maximum charge 3

generate decoy db no

decoy method reverse

decoy prefix DECOY\_

dynamic range 100

peptide minimum length 5



Ввод параметров поиска

New Search

identiprot.theorchromo.ru/data/

matched fragments, min 1

use scoring function hypescore

score threshold 0

show unmatched spectra in results no

report number of sequence candidates 1

[select fixed modifications](#) camC

[select potential modifications](#)

peptide mass shift 0

**Post-search validation**

PSM count

PSMs per protein

charge states

potential modifications

fragment mass tolerance

precursor mass difference

isotopes mass error

missed cleavages

RT difference

Enter search run name

**RUNIDENTIPROT**

Clear chosen files



Послепоисковое  
валидирование  
идентификаций  
(до 9 ортогональных  
критериев)

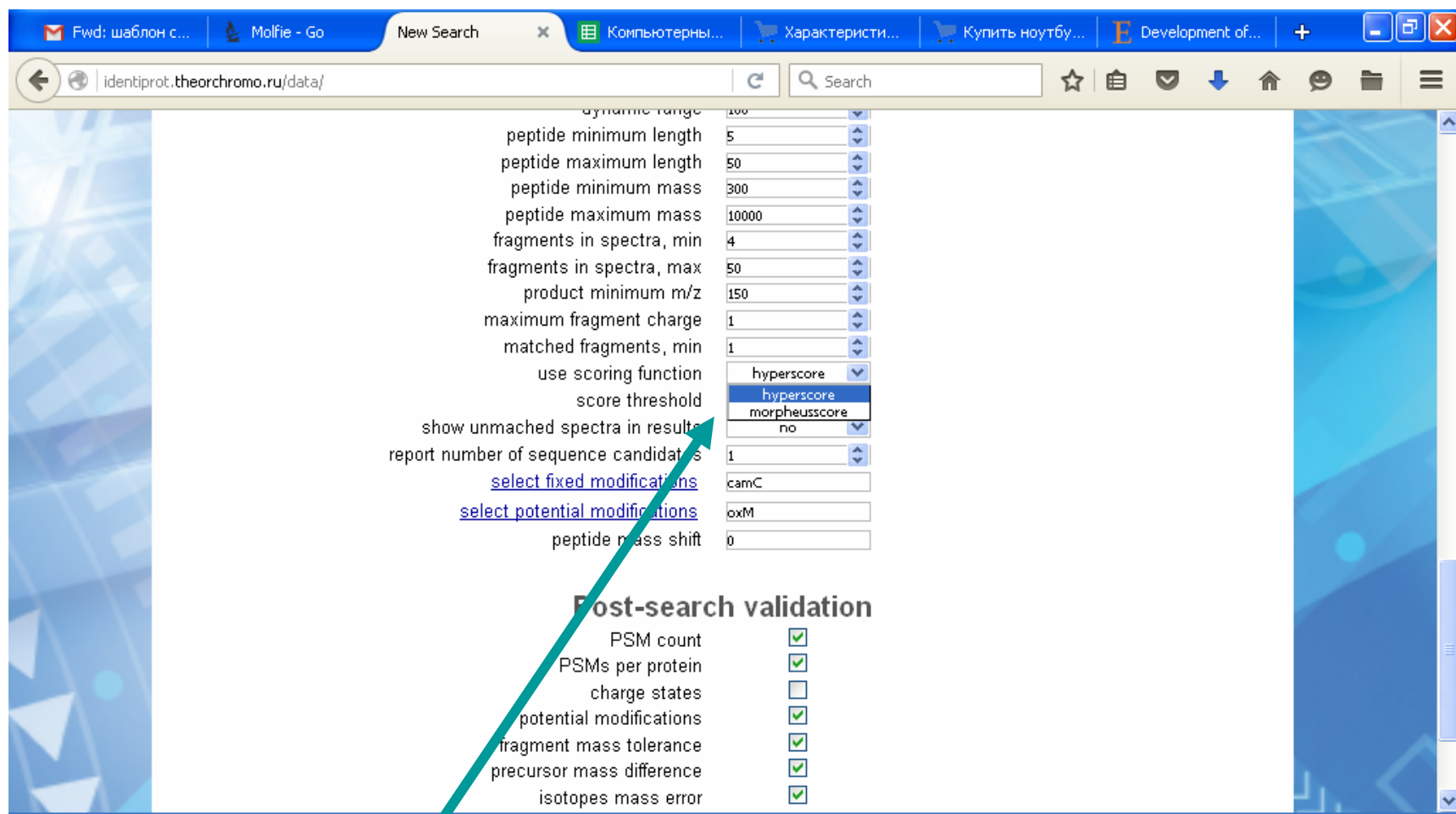
The screenshot shows the IDENTIPROT web interface. At the top, there is a navigation bar with several tabs: "Fwd: шаблон с...", "Molfie - Go", "New Search", "Компьютерны...", "Характеристи...", "Купить ноутбу...", and "Development of...". The address bar shows the URL "identiprot.theorchromo.ru/data/". Below the address bar, there is a search input field and a "Search" button. A blue banner reads "choose preloaded protein database file". Below this is a text input field labeled "Enter search run name" and a large orange button labeled "RUN IDENTIPROT".

The "Specify search parameters" section is active, with three tabs: "MINIMAL", "MEDIUM", and "ADVANCED". The "ADVANCED" tab is selected. The parameters are as follows:

Parameter	Value
send email notification	<input type="checkbox"/>
use auto optimization	<input checked="" type="checkbox"/>
enzyme:	<a href="#">add custom cleavage rule</a> / trypsin
number of missed cleavages	5
precursor accuracy unit	ppm
precursor accuracy left	10
precursor accuracy right	10
precursor isotopic mass error	0
product accuracy, Da	0.05
FDR	1
FDR type	protein
minimum charge	2
maximum charge	5

A green arrow points from the bottom text to the "use auto optimization" checkbox.

**Автоматическая оптимизация входных параметров поиска под конкретные данные**



**Выбор сторонней скоринг-функции в режиме выбора параметров поиска «Advanced»**

The image shows two screenshots of the IdentiPROT web interface. The top screenshot displays the main dashboard with a search history table. A teal arrow points from the 'testsearch01' entry to the second screenshot. The second screenshot shows the search results page for 'testsearch01/test', including a summary table, download options, and three histograms for PSM distributions.

**Вывод результатов поиска**

**START A NEWSEARCH RUN**

**Интерактивный просмотр идентификаций и их распределений по экспериментальным параметрам и физико=химическим свойствам пептидов**

Search name	PSMs	Peptides	Proteins	FDR, %	FDR type	MS/MS	Unfiltered PSMs
test	178	82	32	1.0	psm	3000	2356

Download: [csv files](#) [pepxml files](#) [mgf files](#)  
[figures in png format](#) [figures in svg format](#) [show parameters](#)

**PSM distributions**

- PSMs RT experimental\_min
- PSMs precursor m/z
- PSMs peptide length



**Расчет трех количественных индексов относительного содержания идентифицированных белков**

Search results: 2016-08-10 19-02-45/  
20160809 01 HeLa

Search name	PSMs	Peptides	Proteins	FDR, %	FDR type	MS/MS	Unfiltered PSMs
20160809_01_HeLa	<a href="#">20521</a>	<a href="#">18718</a>	<a href="#">3870</a>	1.0			

Select columns for proteins:

dbname  description  PSMs  peptides  sequence coverage  
 LFQ(SIn)  LFQ(NSAF)  LFQ(emPAI)  protein LN(e value)  all proteins

dbname	PSMs
<a href="#">sp P38646 GRP75_HUMAN</a>	<a href="#">38</a>
<a href="#">sp Q9UKI8-3 ITLK1_HUMAN</a>	<a href="#">1</a>
<a href="#">sp O75694-2 INU155_HUMAN</a>	<a href="#">10</a>
<a href="#">sp P50995 JANX11_HUMAN</a>	<a href="#">8</a>
<a href="#">sp P15880 IRS2_HUMAN</a>	<a href="#">17</a>
<a href="#">sp P07108-4 ACBP_HUMAN</a>	<a href="#">2</a>
<a href="#">sp Q9HC68 CWC22_HUMAN</a>	<a href="#">1</a>

UniProtKB - P38646 (GRP75\_HUMAN)

Protein | Stress-70 protein, mitochondrial  
Gene | HSPA9  
Organism | Homo sapiens (Human)

**Интерактивный режим просмотра белковых identifications**




Fwd: шаблон с... Molfie - Go Contacts x Компьютерны... Характеристики... Купить ноутбу... Development of... +

identiprot.theorchromo.ru/data/ Search

IDENTIPROT START SEARCH UPLOAD FILES SEARCH HISTORY CONTACTS


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**Наши специалисты всегда с вами!**